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       RAW SEQUENCE LISTING              DATE: 08/21/2001
       PATENT APPLICATION:  US/09/671,953  TIME: 11:11:34

       Input Set : A:\Uco991-2.app
       Output Set: N:\CRF3\08162001\I671953.raw
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  3 <110> APPLICANT: Meares, Claude
  4        Chmura, Albert
  5        The Regents of the University of California
  7 <120> TITLE OF INVENTION: Engineering Antibodies That Bind Irreversibly
  9 <130> FILE REFERENCE: 023070-099120US
 11 <140> CURRENT APPLICATION NUMBER: US 09/671,953
 12 <141> CURRENT FILING DATE: 2000-09-27
 14 <150> PRIOR APPLICATION NUMBER: US 60/156,194
 15 <151> PRIOR FILING DATE: 1999-09-27
 17 <150> PRIOR APPLICATION NUMBER: US 60/208,684
 18 <151> PRIOR FILING DATE: 2000-05-31
 20 <160> NUMBER OF SEQ ID NOS: 23
 22 <170> SOFTWARE: PatentIn Ver. 2.1
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ERRORED SEQUENCES

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 443 <211> LENGTH: 360
 444 <212> TYPE: DNA
 445 <213> ORGANISM: Artificial Sequence
 447 <220> FEATURE:
 448 <223> OTHER INFORMATION: Description of Artificial Sequence:V-H sequence of
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 451 <220> FEATURE:
 452 <221> NAME/KEY: CDS
 453 <222> LOCATION: (1)..(360)
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 458  1               5                  10                  15
 460 tcc ctg aaa ctc tcc tgt gca gcc tct gga ttc act tta agt ggt gaa     96
 461 Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Leu Ser Gly Glu
 462             20                  25                  30
 464 acc atg tct tgg gtt cgc cag act ccg gag aag agg ctg gag tgg gtc    144
 465 Thr Met Ser Trp Val Arg Gln Thr Pro Glu Lys Arg Leu Glu Trp Val
 466             35                  40                  45
 468 gca acc act ctt agt ggt ggt ggt ttc acc ttc tat tca gcc agt gtg    192
 469 Ala Thr Thr Leu Ser Gly Gly Gly Phe Thr Phe Tyr Ser Ala Ser Val
 470         50                  55                  60
 472 aag ggt cgt ttc acc atc tcc aga gac aat gcc cag aac aac ctc tat    240
 473 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Gln Asn Asn Leu Tyr
 474  65                  70                  75                  80
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476 cta caa ctg aat agt ctg agg tct gag gac acg gcc ttg tat ttc tgt   288
477 Leu Gln Leu Asn Ser Leu Arg Ser Glu Asp Thr Ala Leu Tyr Phe Cys
478             85                  90                  95
480 gca agt cat cgg ttt gtt cac tgg ggc cac ggg act ctg gtc act gtc   336
481 Ala Ser His Arg Phe Val His Trp Gly His Gly Thr Leu Val Thr Val
482            100                 105                 110
484 tct gca gcc aaa acg aca ccc cca                                    360
485 Ser Ala Ala Lys Thr Thr Pro Pro
486            115                 120
489 <210> SEQ ID NO: 17
490 <211> LENGTH: 120
491 <212> TYPE: PRT
492 <213> ORGANISM: Artificial Sequence
493 <220> FEATURE:                                                     W-->
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496 <211> LENGTH:                                                      W-->
496 <212> TYPE:                                                        W-->
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500 Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Leu Ser Gly Glu
501            20                  25                  30
503 Thr Met Ser Trp Val Arg Gln Thr Pro Glu Lys Arg Leu Glu Trp Val
504            35                  40                  45
506 Ala Thr Thr Leu Ser Gly Gly Gly Phe Thr Phe Tyr Ser Ala Ser Val
507         50                  55                  60
509 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Gln Asn Asn Leu Tyr
510  65                 70                  75                  80
512 Leu Gln Leu Asn Ser Leu Arg Ser Glu Asp Thr Ala Leu Tyr Phe Cys
513            85                  90                  95
515 Ala Ser His Arg Phe Val His Trp Gly His Gly Thr Leu Val Thr Val
516            100                 105                 110
518 Ser Ala Ala Lys Thr Thr Pro Pro
519            115                 120
```

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/671,953

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Input Set : **A:\Uco991-2.app**
Output Set: **N:\CRF3\08162001\I671953.raw**

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L:172 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:230 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:289 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:449 M:254 E: No. of Bases conflict, LENGTH:Input:255 Counted:3 SEQ:16
L:449 M:112 C: (48) String data converted to lower case,
L:449 M:252 E: No. of Seq. differs, <211>LENGTH:Input:360 Found:3 SEQ:16
L:455 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:455 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:455 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:455 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:493 M:258 W: Mandatory Feature missing, <220> FEATURE:
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L:496 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:496 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:496 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:496 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:600 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:609 M:258 W: Mandatory Feature missing, <220> FEATURE:
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